

**Nucleotides 1-1560 of the genome of
Enterovirus type 71 BrCr (GenBank Accession No.: U22521)**

1 TTA AACAGC TGTGGGTTGT CACCCACCCA CAGGGTCCAC TGGGCGCTAG TACACTGGTA
61 TCTCGGTACC TTTGTACGCC TGTTTTATAC CCCCTCCCTG ATTTGCAACT TAGAAGCAAC
 f1 (72-87)
121 GCAAACCAGA TCAATAGTAG GTGTGACATA CCAGTCCCAT CTTGATCAAG CACTTCTGTA
181 TCCCCGGGACC GAGTATCAAT AGACTGTGCA CACGGTTGAA GGAGAAAACG TCCGTTACCC
 f2 (167-187)
241 GGCTAACCTAC TTCGAGAAGC CTAGTAACGC CATTGAAGTT GCAGAGTGTT TCGCTCAGCA
 f3 (248-266)
301 CTCCCCCGGT GTAGATCAGG TCGATGAGTC ACCGCATTCC CCACGGGCGA CCGTGGCGGT
361 GGCTGCGTTG GCGGCCTGCC TATGGGGTAA CCCATAGGAC GCTCTAATAC GGACATGGCG
421 TGAAGAGTCT ATTGAGCTAG TTAGTAGTCC TCCGCCCCCT GAATGCGGCT AATCCTAACT
 f5 (423-439) **p1 (448-474)**
481 GCGGAGCACA TACCCTTAAT CCAAAGGGCA GTGTGTCGTA ACGGGCAACT CTGCAGCGGA
 p2 (514-546)
541 ACCGACTACT TTGGGTGTCC GTGTTTCTTT TTAATCTTGT ATTGGCTGCT TATGGTGACA
 p3 (547-574) **r2 (583-602)**
601 ATTAAGAAT TGTACCATA TAGCTATTGG ATTGGCCATC CAGTGTCAAA CAGAGCTATT
 r2 (583-602) **r1 (627-645)**
661 GTATATCTCT TTGTTGGATT CACACCTCTC ACTCTTGAAA CGTTACACAC CCTCAATTAC
721 ATTATACTGC TGAACACGAA GCGATGGGCT CCCAGGTCTC CACACAGCGA TCCGGCTCGC
781 ATGAGAATTC CAACTCAGCC ACGGAAGGCT CCACTATAAA TTACACAACC ATTAATTACT
841 ACAAGAGACT GTATGCTGCC ACTGCTGGAA AGCAAAGTCT CAAACAAGAT CCTGACAAGT
901 TTGCGAACCCT TGTGAAGGAC ATCTTTACTG AAATGGCAGC GCCCTTAAAG TCTCCCTCTG
961 CTGAAGCATG TGGCTATAGC GACCGAGTGG CACAGCTTAC CATTGGAAAT TCCACCATTA
1021 CTACACAAGA AGCAGCAAAC ATAATAGTTG GGTATGGTGA GTGGCCTTCA TACTGCTCTG
1081 ATAATGATGC AACAGCGGTA GACAAACCTA CACGGCCTGA TGTCTCAGTA AATAGATTTT
1141 ACACGCTAGA CACTAAGCTA TGGGAGAAAT CATCCAAGGG GTGGTACTGG AAGTTCCAG
 f7 (1179-1198)
1201 ATGTA CTGAC TGAAACCGGA GTTTTGGTGC CAAATGCACA ATTTCACTAC TTATACCGTT
1261 CAGGGTTCTG CATCCACGTT CAATGTAACG CTAGCAAATT TCACCAAGGG GCGCTACTCG
1321 TTGCGGTATT GCCCGAGTAT GTCATTGGAA CAGTGGCAGG CGGCACAGGC ACAGAGAACA
 16-1 (1343-1373)
1381 GTCACCCTCC TTATAACAA ACCCAACCCG GCGCTGATGG ATTTGAATTA CAACATCCAT
 71-2 (1390-1419)/16-2 (1390-1419)
1441 ATGTTCTTGA TGCTGGAATT CCAATATCTC AGTTGACAGT GTGCCTCAC CAGTGGATCA
 71-3 (1453-1481) **r4 (1485-1504)**
1501 ATTTACGAAC CAACAATTGT GCCACCATAA TAGTGCCATA CATGAACACA CTACCTTTTG

Figure 1